

SEQUENCE LISTING

<110> HOSHINO, Tatsuo
OJIMA, Kazuyuki
SETOGUCHI, Yutaka

<120> ASTAXANTHIN SYNTHETASE

<130> ASTAXANTHIN SYNTHETASE

<140> US/09/518,386

<141> 2000-03-03

<150> EP 99104668.1

<151> 1999-03-09

<150> EP 00101666.6

<151> 2000-02-01

<160> 32

<170> PatentIn Ver. 2.1

<210> 1

<211> 557

<212> PRT

<213> Phaffia rhodozyma

<220>

<221> TRANSIT

<222> (1)..(26)

<400> 1

Met	Phe	Ile	Leu	Val	Leu	Leu	Thr	Gly	Ala	Leu	Gly	Leu	Ala	Ala	Phe
1					5				10					15	

Ser	Trp	Ala	Ser	Ile	Ala	Phe	Phe	Ser	Leu	Tyr	Leu	Ala	Pro	Arg	Arg
			20					25					30		

Ser	Ser	Leu	Tyr	Asn	Leu	Gln	Gly	Pro	Asn	His	Thr	Asn	Tyr	Phe	Thr
		35					40						45		

Gly Asn Phe Leu Asp Ile Leu Ser Ala Arg Thr Gly Glu Glu His Ala
 50 55 60

Lys Tyr Arg Glu Lys Tyr Gly Ser Thr Leu Arg Phe Ala Gly Ile Ala
 65 70 75 80

Gly Ala Pro Val Leu Asn Ser Thr Asp Pro Lys Val Phe Asn His Val
 85 90 95

Met Lys Glu Ala Tyr Asp Tyr Pro Lys Pro Gly Met Ala Ala Arg Val
 100 105 110

Leu Arg Ile Ala Thr Gly Asp Gly Val Val Thr Ala Glu Gly Glu Ala
 115 120 125

His Lys Arg His Arg Arg Ile Met Ile Pro Ser Leu Ser Ala Gln Ala
 130 135 140

Val Lys Ser Met Val Pro Ile Phe Leu Glu Lys Gly Met Glu Leu Val
 145 150 155 160

Asp Lys Met Met Glu Asp Ala Ala Glu Lys Asp Met Ala Val Gly Glu
 165 170 175

Ser Ala Gly Glu Lys Lys Ala Thr Arg Leu Glu Thr Glu Gly Val Asp
 180 185 190

Val Lys Asp Trp Val Gly Arg Ala Thr Leu Asp Val Met Ala Leu Ala
 195 200 205

Gly Phe Asp Tyr Lys Ser Asp Ser Leu Gln Asn Lys Thr Asn Glu Leu
 210 215 220

Tyr Val Ala Phe Val Gly Leu Thr Asp Gly Phe Ala Pro Thr Leu Asp
 225 230 235 240

Ser Phe Lys Ala Ile Met Trp Asp Phe Val Pro Tyr Phe Arg Thr Met
 245 250 255

Lys Arg Arg His Glu Ile Pro Leu Thr Gln Gly Leu Ala Val Ser Arg
 260 265 270

Arg Val Gly Ile Glu Leu Met Glu Gln Lys Lys Gln Ala Val Leu Gly

10066607.020402

10065007.030403

275		280		285
Ser Ala Ser Asp Gln Ala Val Asp Lys Lys Asp Val Gln Gly Arg Asp				
290		295		300
Ile Leu Ser Leu Leu Val Arg Ala Asn Ile Ala Ala Asn Leu Pro Glu				
305		310		315 320
Ser Gln Lys Leu Ser Asp Glu Glu Val Leu Ala Gln Ile Ser Asn Leu				
		325		330 335
Leu Phe Ala Gly Tyr Glu Thr Ser Ser Thr Val Leu Thr Trp Met Phe				
		340		345 350
His Arg Leu Ser Glu Asp Lys Ala Val Gln Asp Lys Leu Arg Glu Glu				
		355		360 365
Ile Cys Gln Ile Asp Thr Asp Met Pro Thr Leu Asp Glu Leu Asn Ala				
		370		375 380
Leu Pro Tyr Leu Glu Ala Phe Val Lys Glu Ser Leu Arg Leu Asp Pro				
385		390		395 400
Pro Ser Pro Tyr Ala Asn Arg Glu Cys Leu Lys Asp Glu Asp Phe Ile				
		405		410 415
Pro Leu Ala Glu Pro Val Ile Gly Arg Asp Gly Ser Val Ile Asn Glu				
		420		425 430
Val Arg Ile Thr Lys Gly Thr Met Val Met Leu Pro Leu Phe Asn Ile				
		435		440 445
Asn Arg Ser Lys Phe Ile Tyr Gly Glu Asp Ala Glu Glu Phe Arg Pro				
		450		455 460
Glu Arg Trp Leu Glu Asp Val Thr Asp Ser Leu Asn Ser Ile Glu Ala				
465		470		475 480
Pro Tyr Gly His Gln Ala Ser Phe Ile Ser Gly Pro Arg Ala Cys Phe				
		485		490 495
Gly Trp Arg Phe Ala Val Ala Glu Met Lys Ala Phe Leu Phe Val Thr				
		500		505 510

Leu Arg Arg Val Gln Phe Glu Pro Ile Ile Ser His Pro Glu Tyr Glu
 515 520 525

His Ile Thr Leu Ile Ile Ser Arg Pro Arg Ile Val Gly Arg Glu Lys
 530 535 540

Glu Gly Tyr Gln Met Arg Leu Gln Val Lys Pro Val Glu
 545 550 555

<210> 2
 <211> 1932
 <212> DNA
 <213> Phaffia rhodozyma

<220>
 <221> CDS
 <222> (33)..(1706)

<220>
 <221> polyA_site
 <222> (1871)

<220>
 <221> mRNA
 <222> (14)..(1891)

<400> 2
 gaattcggca cgaggccacc tactttctcc at atg ttc atc ttg gtc ttg ctc
 53

Met Phe Ile Leu Val Leu Leu
 1 5

aca ggt gct tta ggc ctg gct gct ttc tca tgg gca tcc ata gcg ttc
 101

Thr Gly Ala Leu Gly Leu Ala Ala Phe Ser Trp Ala Ser Ile Ala Phe
 10 15 20

ttc agt ctt tac ctc gct ccg agg cga tct tca ctg tat aac ctt cag
 149

Phe Ser Leu Tyr Leu Ala Pro Arg Arg Ser Ser Leu Tyr Asn Leu Gln
 25 30 35

1006607.030100

gca aac atc gcc gcc aac ctg cct gaa tct caa aag ctg tcc gat gag
1013

Ala Asn Ile Ala Ala Asn Leu Pro Glu Ser Gln Lys Leu Ser Asp Glu
315 320 325

gag gta ctc gct cag atc agt aac ctg tta ttt gct gga tat gaa act
1061

Glu Val Leu Ala Gln Ile Ser Asn Leu Leu Phe Ala Gly Tyr Glu Thr
330 335 340

tct tcg aca gtc ttg aca tgg atg ttt cac cga ctc tca gaa gac aaa
1109

Ser Ser Thr Val Leu Thr Trp Met Phe His Arg Leu Ser Glu Asp Lys
345 350 355

gcc gtt cag gat aaa ctt cga gaa gaa att tgt cag atc gac acg gat
1157

Ala Val Gln Asp Lys Leu Arg Glu Glu Ile Cys Gln Ile Asp Thr Asp
360 365 370 375

atg cct acg cta gac gaa ctt aat gcg ttg cct tat ctc gaa gcg ttt
1205

Met Pro Thr Leu Asp Glu Leu Asn Ala Leu Pro Tyr Leu Glu Ala Phe
380 385 390

gtt aag gag tct ctt cgt cta gac cct cct agt ccg tat gct aac cgt
1253

Val Lys Glu Ser Leu Arg Leu Asp Pro Pro Ser Pro Tyr Ala Asn Arg
395 400 405

gaa tgc tta aag gat gaa gac ttc atc cca ctt gcc gag cct gtc att
1301

Glu Cys Leu Lys Asp Glu Asp Phe Ile Pro Leu Ala Glu Pro Val Ile
410 415 420

ggc cga gat ggg tcg gtc atc aac gag gtc cgg atc acg aaa gga acg
1349

Gly Arg Asp Gly Ser Val Ile Asn Glu Val Arg Ile Thr Lys Gly Thr
425 430 435

atg gtc atg ctt ccg ttg ttc aac atc aat cgt tca aag ttc att tat
1397

Met Val Met Leu Pro Leu Phe Asn Ile Asn Arg Ser Lys Phe Ile Tyr

440

445

450

455

gga gaa gat gca gaa gaa ttc aga ccg gag agg tgg ctt gag gac gta
1445

Gly Glu Asp Ala Glu Glu Phe Arg Pro Glu Arg Trp Leu Glu Asp Val
460 465 470

aca gac tcg ctc aac agt att gaa gca ccc tat gga cac cag gcg agc
1493

Thr Asp Ser Leu Asn Ser Ile Glu Ala Pro Tyr Gly His Gln Ala Ser
475 480 485

ttt atc tct gga ccc aga gct tgc ttt ggt tgg cga ttt gct gtc gcc
1541

Phe Ile Ser Gly Pro Arg Ala Cys Phe Gly Trp Arg Phe Ala Val Ala
490 495 500

gag atg aag gcc ttc ttg ttt gtc act ctc cgt cgg gtc cag ttc gag
1589

Glu Met Lys Ala Phe Leu Phe Val Thr Leu Arg Arg Val Gln Phe Glu
505 510 515

ccc atc atc tct cat cca gag tac gag cac atc acc ttg atc att tcc
1637

Pro Ile Ile Ser His Pro Glu Tyr Glu His Ile Thr Leu Ile Ile Ser
520 525 530 535

cgt cct cga atc gtt ggt aga gag aag gag ggg tac cag atg cgt ttg
1685

Arg Pro Arg Ile Val Gly Arg Glu Lys Glu Gly Tyr Gln Met Arg Leu
540 545 550

cag gtc aag ccg gtc gaa tga gttgattcct catatgttaa gagaagttct
1736

Gln Val Lys Pro Val Glu
555

atatctgaga atgtgtgact aggacaatgc cttctttgta tcgatttggt tctcataccc
1796

gggcaggcgc tatgacttct acgtcgtcta tcgtcgtctt ggactctctt cttaccctat
1856

1005607-030403

atattattcc atccgaaaaa aaaaaaaaaa aaaaaaaaaa aaaaagcggc cgctcgagcc
1916

ggctcgtgcc gaattc
1932

<210> 3
<211> 557
<212> PRT
<213> Phaffia rhodozyma

<400> 3
Met Phe Ile Leu Val Leu Leu Thr Gly Ala Leu Gly Leu Ala Ala Phe
1 5 10 15
Ser Trp Ala Ser Ile Ala Phe Phe Ser Leu Tyr Leu Ala Pro Arg Arg
20 25 30
Ser Ser Leu Tyr Asn Leu Gln Gly Pro Asn His Thr Asn Tyr Phe Thr
35 40 45
Gly Asn Phe Leu Asp Ile Leu Ser Ala Arg Thr Gly Glu Glu His Ala
50 55 60
Lys Tyr Arg Glu Lys Tyr Gly Ser Thr Leu Arg Phe Ala Gly Ile Ala
65 70 75 80
Gly Ala Pro Val Leu Asn Ser Thr Asp Pro Lys Val Phe Asn His Val
85 90 95
Met Lys Glu Ala Tyr Asp Tyr Pro Lys Pro Gly Met Ala Ala Arg Val
100 105 110
Leu Arg Ile Ala Thr Gly Asp Gly Val Val Thr Ala Glu Gly Glu Ala
115 120 125
His Lys Arg His Arg Arg Ile Met Ile Pro Ser Leu Ser Ala Gln Ala
130 135 140
Val Lys Ser Met Val Pro Ile Phe Leu Glu Lys Gly Met Glu Leu Val
145 150 155 160
Asp Lys Met Met Glu Asp Ala Ala Glu Lys Asp Met Ala Val Gly Glu
165 170 175
Ser Ala Gly Glu Lys Lys Ala Thr Arg Leu Glu Thr Glu Gly Val Asp
180 185 190
Val Lys Asp Trp Val Gly Arg Ala Thr Leu Asp Val Met Ala Leu Ala
195 200 205
Gly Phe Asp Tyr Lys Ser Asp Ser Leu Gln Asn Lys Thr Asn Glu Leu
210 215 220
Tyr Val Ala Phe Val Gly Leu Thr Asp Gly Phe Ala Pro Thr Leu Asp
225 230 235 240

<210> 4
<211> 3969
<212> DNA
<213> Phaffia rhodozyma

<220>
<221> 5'UTR
<222> (517)..(518)

<220>
<221> intron
<222> (784)..(898)

<220>
<221> intron
<222> (1016)..(1087)

<220>
<221> intron
<222> (1180)..(1302)

<220>
<221> intron
<222> (1518)..(1600)

<220>
<221> intron
<222> (1635)..(1723)

<220>
<221> intron
<222> (1867)..(1939)

<220>
<221> intron
<222> (2000)..(2081)

<220>
<221> intron
<222> (2182)..(2257)

<220>
<221> intron

J00607-00106

$\langle 222 \rangle \quad (2355) \dots (2431)$ $\langle 220 \rangle$ $\langle 220 \rangle$ $\langle 220 \rangle$

$\langle 220 \rangle$

<220>

<220>

<220>

 $\langle 220 \rangle$ $\langle 220 \rangle$

<400> 4

tggaggagcg ggagcgggag gagcagcagg tgatgcatcg ggtggacaga atcagtagtg
120

1440
tggccgtggg agagtcggcc ggtgaaaaga aggcaaccag actcgagacc gaaggagtcg
1500
atgtaaagga ttgggtcgtg agtaccgcgc tattccttca ccttgatgga cgaagcatat
1560
caaggaaagg ttcattgact gacaaacact atcttaccag ggtcgagcta ctctggacgt
1620
catggctctt gcaggtcagt ctactctctc ttataaatgc tccacatatg tatgcatgta
1680
ctgacatgct cttcctatat tcgatacgac gtcatatgtc caggatttga ctataagagc
1740
gactcgctcc agaacaagac caatgagctc tatgtcgctt ttgtcggact taccgatggg
1800
tttgtccta ccttggactc gttcaaggct atcatgtggg attttgtacc ttacttccga
1860
actatggtat gtctgccatt ctttgatata caaagattat ggataggtta cttgctaaaa
1920
tttcacctat cgtgaacaga aacggagaca tgagatacct ttgactcaag gattagcagt
1980
ttcccgacga gttgggatcg taagtgccag atcaagcctc tctgaatatt cttgggtcatc
2040
atcttaacct cctaggctca ttcattccatg gtgcgcaata ggagcttatg gagcaaaaga
2100
agcaggccgt gcttggctca gcttccgatc aggctgttga taaaaggat gttcaaggtc
2160
gggatatact aagtctccta ggtagtaac gtttttaaac gtatatacag agcggcgaca
2220
ttctttccct gacaactgtc aacatgctcg ttactagtga gagcaaacat cgccgccaac
2280
ctgcctgaat ctcaaaagct gtccgatgag gaggtactcg ctcagatcag taacctgtta
2340
tttgctggat atgagtgtgt atcctttccc ctctctatcc ttagctgatt aaaagcacta
2400
atagaggtct ttatgtttcc tgtttgatca gaacttcttc gacagtcttg acatggatgt
2460
ttcaccgact ctcaagaagac aaagccgttc aggataaact tcgagaagaa atttgtcaga
2520
tcgacacgga tatgcctacg ctgtgaggat gtttttgatg ctaaattact tcttcttgca
2580
aatgactaaa acggccttcc attcttgatc catttttagag acgaacttaa tgcgttgcc
2640
tatctcgaag cggttggttc tcgattcttg gtcttgtctt ccaaatacaa tacggattat
2700

tgctcatctg atttgcgtct acgggctgtg gaatttaact agtttggttaa ggagtctctt
 2760
 cgtctagacc ctccctagtcc gtatgctaac cgtgaatgct taaaggatga agacgtatgt
 2820
 tggcttcac acgcataatt ttcatttcat attcctttgt acatacgcac acaggctgac
 2880
 cgagctcaaa ttccggcttc ctcttctgtg cttctttttc tggcctttct tatcttcatt
 2940
 cttcaaccaa aatttgtcac agttcatccc acttgccgag cctgtcattg gtcgagatgg
 3000
 gtcggtcac aacgaggtcc ggatcacgaa aggaacgatg gtcatgcttc gtaagttttc
 3060
 ctttatttca tctcgtccat gaaatagttt ctgatagacg cggaccaatt cagcgttggt
 3120
 caacatcaat cgttcaaagt tcatttatgg agaagatgca gaagaattca ggtacaattc
 3180
 gttttctttt aaaagccaat cggtttcgta tcgtaattga ccgggctctc ttttaatttc
 3240
 tcgaaagacc ggagaggtgg cttgaggacg taacagactc gctcaacagt attgaagcac
 3300
 cctatggaca ccaggcgagc tgtatgtttt attgatttta tctttgtgaa ttttgcaaaa
 3360
 cgttgaactt cgcgcttccc ttgttggtga aatcccagtt atctctggac ccagagcttg
 3420
 cttgtaagtt tcttctcatc tggcgcctta gcagtatccg atcagccatc tagttctttg
 3480
 tacgattgtt tctgactctc tcgactttcg cagtgggttg cgatttgctg tcgccgagat
 3540
 gaaggccttc ttgtttgtca ctctccgtcg ggtccagttc gagcccatca tctctcatcc
 3600
 agagtacgag cacatcacct tgatcatttc ccgtcctoga atcgttggta gagagaagga
 3660
 ggggtaccag atgcgtttgc aggtcaagcc ggtcgaatga gttgattctt catatgttaa
 3720
 gagaagttct atatctgaga atgtgtgact aggacaatgc cttctttgta tcgatttggt
 3780
 tctcatacc gccgcaggcgc tatgacttct acgtcgtcta tcgtcgtctt ggactctctt
 3840
 cttaccctat atattattcc atccgtctgt atatttgtct atcacgacgt ctgtgtcgtc
 3900
 aactcaatat tcagcctctt catgcttctg tgtctccata gatgtgatct tcatgtttgt
 3960
 cgactgcag

3969

<210> 5
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Sense primer
for expression of the AST gene in E. coli

<400> 5
gttcaaagtt catttatgga
20

<210> 6
<211> 47
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Antisense
primer for expression of the AST gene in E. coli

<400> 6
ggatcctcag tgggtggtggt ggtggtgttc gaccggcttg acctgca
47

<210> 7
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 5' sense
primer for expression of a modified AST gene in E.
coli

<400> 7
catatgcacc accaccacca ccacctgtat aaccttcagg ggccc

<210> 8
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 5' antisense
primer for expression of a modified AST gene in E.
coli

<400> 8
gtaacaacac catctccggt
20

<210> 9
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 3' anti sense
primer for expression of a modified AST gene in E.
coli

<400> 9
ggatcctcaa ctcattcgac cggctt
26

<210> 10
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Genome walking
primer for cloning of the AST gene

<400> 10

tagagagaag gaggggtacc agatgc
26

<210> 11
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Antisense
primer for cloning of the terminator region of the
AST gene

<400> 11
ccccggattg tggagaaact
20

<210> 12
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Sense primer
for cloning the genomic AST gene

<400> 12
atgttcatct tggctcttgct
20

<210> 13
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Antisense
primer for cloning the genomic AST gene

<400> 13

acgtagaagt catagcgcct
20

<210> 14
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Sense primer
for RT-PCR of the AST gene

<400> 14
tttgactcaa ggattagcag
20

<210> 15
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Antisense
primer for RT-PCR of the AST gene

<400> 15
tgtcttctga gagtcggtga
20

<210> 16
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Degenerate
sense primer for cloning of the TPI gene

<220>
<221> misc_feature

1006607.634103

<222> 3
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 6
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 15
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 18
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 21
<223> n is a or c or g or t

<400> 16
mgnacnttyt tygtnggngg naay
24

<210> 17
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Degenerate
antisense primer for cloning the TPI gene

<220>
<221> misc_feature
<222> 3
<223> n is a or c or g or t

<220>

```
<221> misc_feature
<222> 6
<223> n is a or c or g or t
```

<210> 19
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Nested walking
primer for cloning of the TPI terminator

<400> 19
ggatctgtct ctgcctccaa ctgcaag
27

<210> 20
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primary
walking primer for cloning of the TPI promoter

<400> 20
gggtcaatgt cggcagcgag aagccca
27

<210> 21
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Nested walking
primer for cloning of the TPI promoter

<400> 21
atgtactcgg tagcactgat caagtag
27

<210> 22

<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Sense primer
for construction of the TPI promoter cassette

<400> 22
gcggccgcat ccgtctcgcc atcagtct
28

<210> 23
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Antisense
primer for construction of the TPI promoter
cassette

<400> 23
cctgcaggtc tagagatgaa taaatataaa gagt
34

<210> 24
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Sense primer
for construction of the TPI terminator cassette

<400> 24
cctgcaggta aatatatcca gggattaacc ccta
34

<210> 25

10056007.030103

<211> 28
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Antisense
primer for construction of the TPI terminator
cassette

<400> 25
ggtacccgtg cgcagtcgac cgagacat
28

<210> 26
<211> 35
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Degenerate
sense primer for cloning of the AMY gene

<220>
<221> misc_feature
<222> 15
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 21
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 27
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 30
<223> n is a or c or g or t

<400> 26
gaytayathc arggnatggg nttyrmngcn athtg
35

<210> 27
<211> 35
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Degenerate
antisense primer for cloning of the AMY gene

<220>
<221> misc_feature
<222> 6
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 9
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 24
<223> n is a or c or g or t

<220>
<221> misc_feature
<222> 30
<223> n is a or c or g or t

<400> 27
tgytcngtnc crtartadat datnggdatn ccrtc
35

<210> 28
<211> 26
<212> DNA
<213> Artificial Sequence

10066007 DE0102

<220>

<223> Description of Artificial Sequence: Sense primer
for construction of a partial AMY cassette

<400> 28

ccgcggcatt gatacctcta ccccg

26

<210> 29

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Antisense
primer for construction of a partial AMY cassette

<400> 29

gcggccgcct gcaatcctgg atccaccg

28

<210> 30

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Sense primer
for construction of the AST cassette

<400> 30

tctagaatgt tcattcttggt cttgctca

28

<210> 31

<211> 28

<212> DNA

<213> Artificial Sequence

Figure 1 displays 12 ECG strips, each showing a different cardiac rhythm. The strips are labeled with patient names and clinical details. The rhythms shown include normal sinus rhythm, sinus tachycardia, sinus bradycardia, sinus arrhythmia, sinus pause, sinus arrest, sinus exit block, and sinus exit block with compensatory pause.

```
<400> 31
cctgcaggtc attcgaccgg cttgacct
28
```

<220>
<223> Description of Artificial Sequence: Sense primer
for confirmation of integration at the AMY locus
by PCR analysis

Page 27